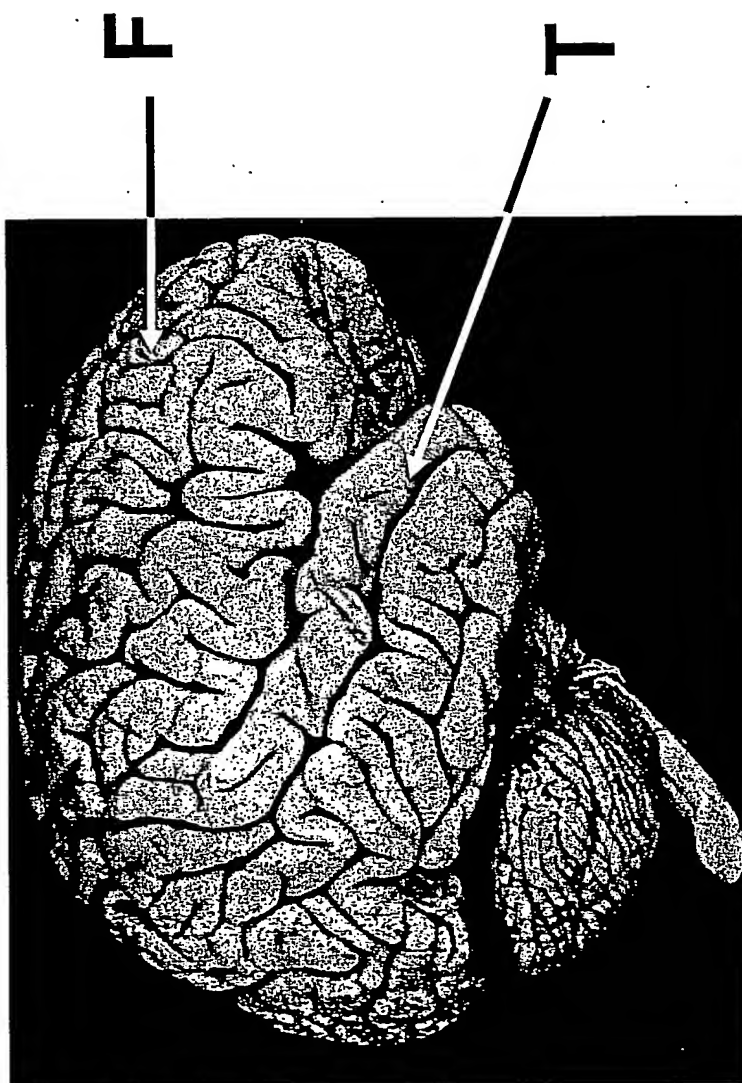


Fig. 1: Identification of Genes Involved
in Alzheimer's Disease Pathology



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Fig. 2: Identification of differentially expressed genes in a suppressive subtractive hybridization screen by dot blot analysis

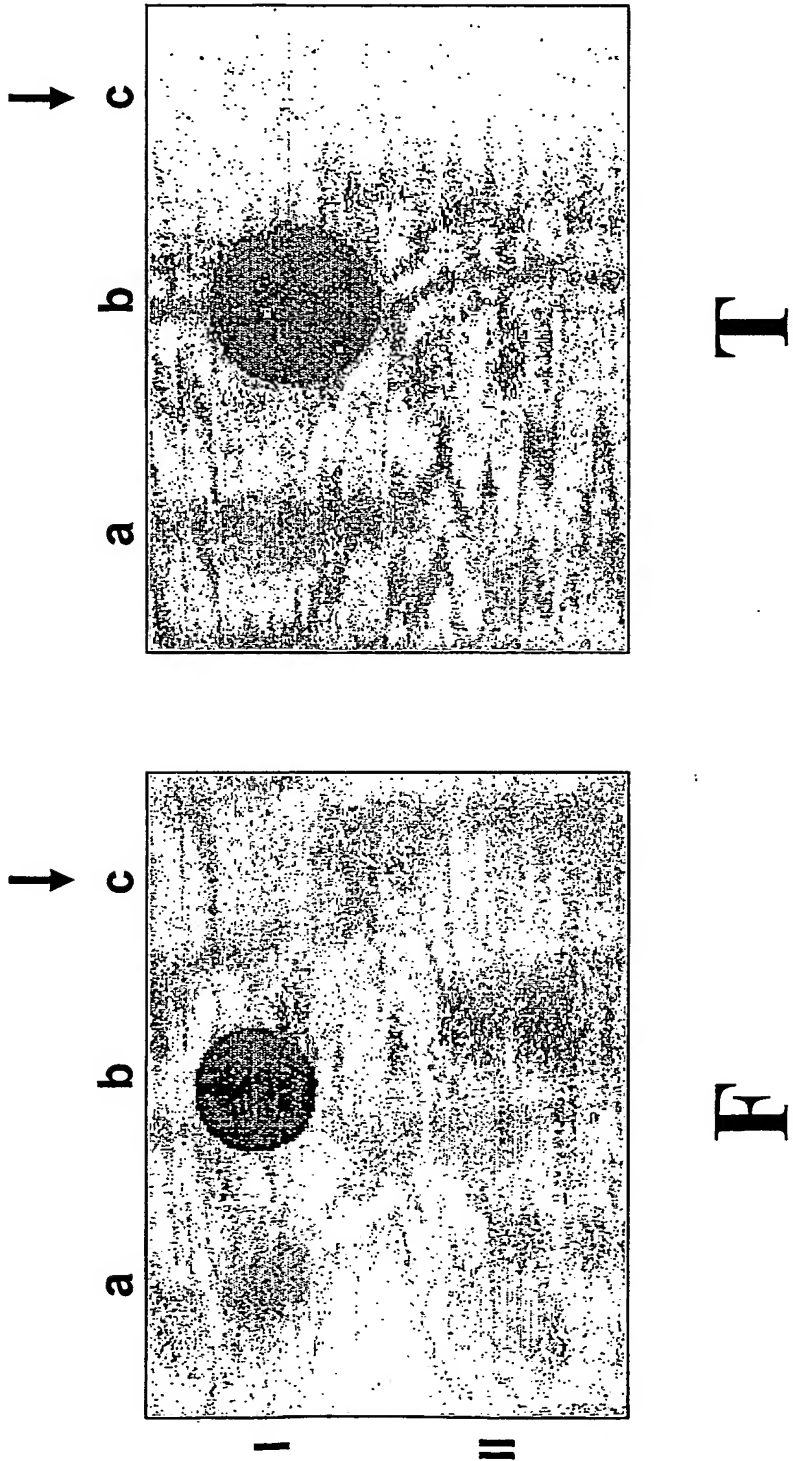


Fig. 3: Verification of differential expression of SCN2A by quantitative RT-PCR

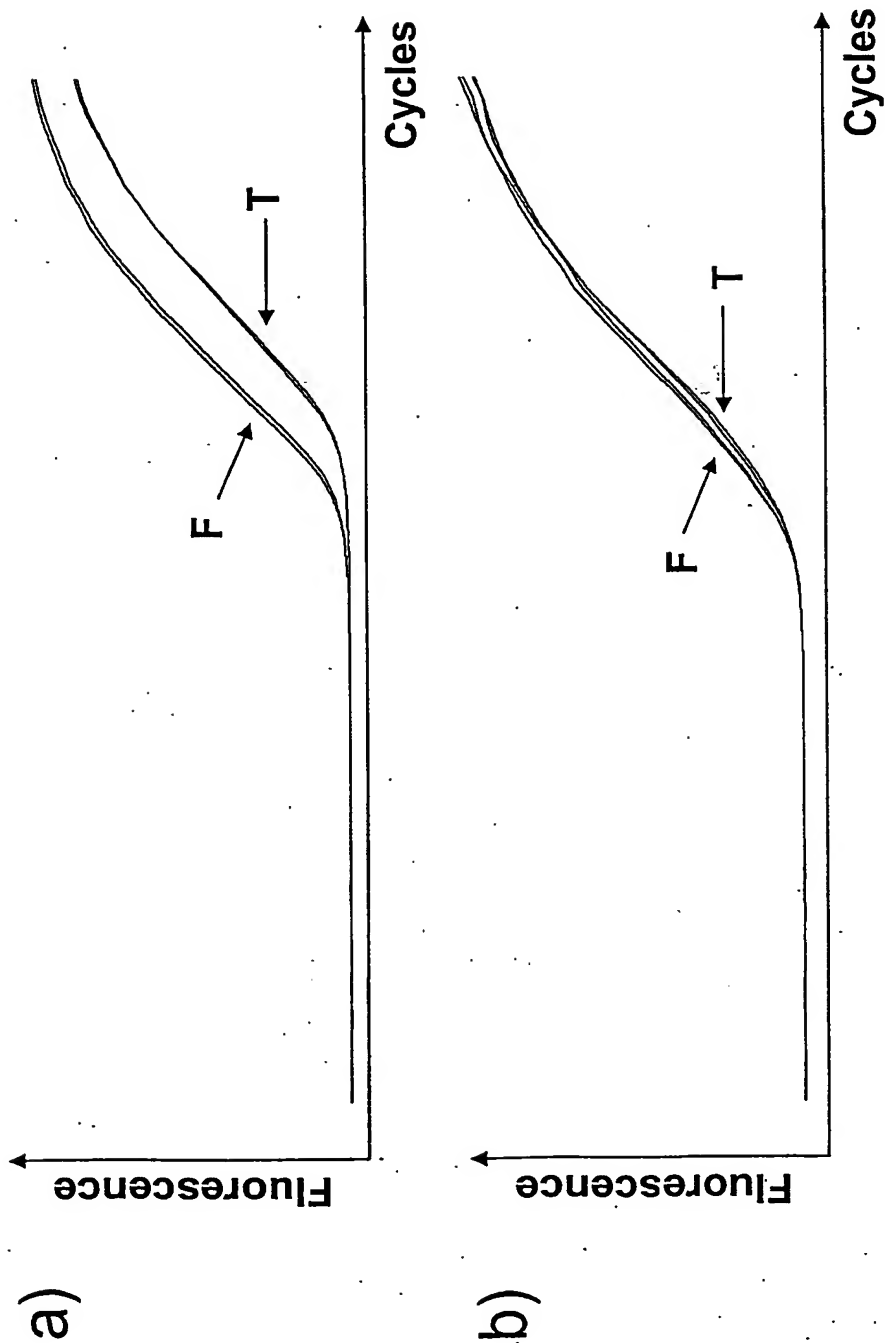
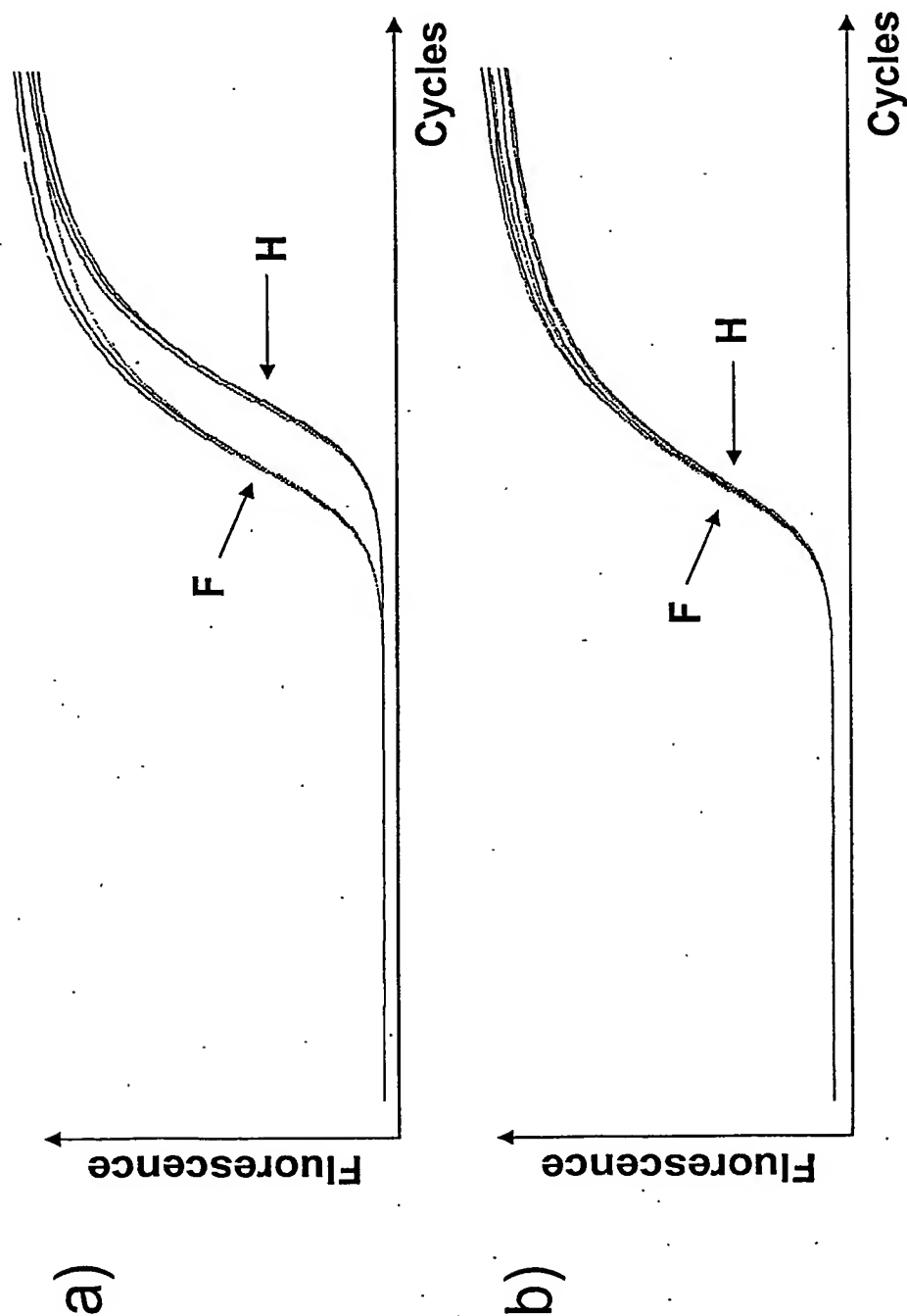


Fig. 4: Verification of differential expression of SCN2A by quantitative RT-PCR



**Fig. 5: Nucleotide sequence
of SEQ ID NO: 1**

Length: 272 bp

```
1  AATTAAGGTT GGAAGAATAA AAAGCAAGAA GCTCTTCCTT GTTTGCTGCA
51  ACCTATTGCT TAATGACATG AAGAATGAGG TCTTGGTAGA ACAATTTGCT
101 TCACTTTACC ACTGATATAT GGCTTCCCAT ATTAGACTTC TGAACAGGGG
151 AAGGAATAAG ATACAGCAGC ATAGGCAAGA TAAACATGCA GCAGTGACAG
201 CTTCAAAC TAATGGAACC AATTACATCA TATTACCTGT TGGAAGCTTG
251 CAAACTATAC TTACTGGGGT AC
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**Fig. 6: Schematic alignment of SEQ ID NO: 1
with sodium channel type II alpha subunit
(Accession No. AF327224 - AF327246)**

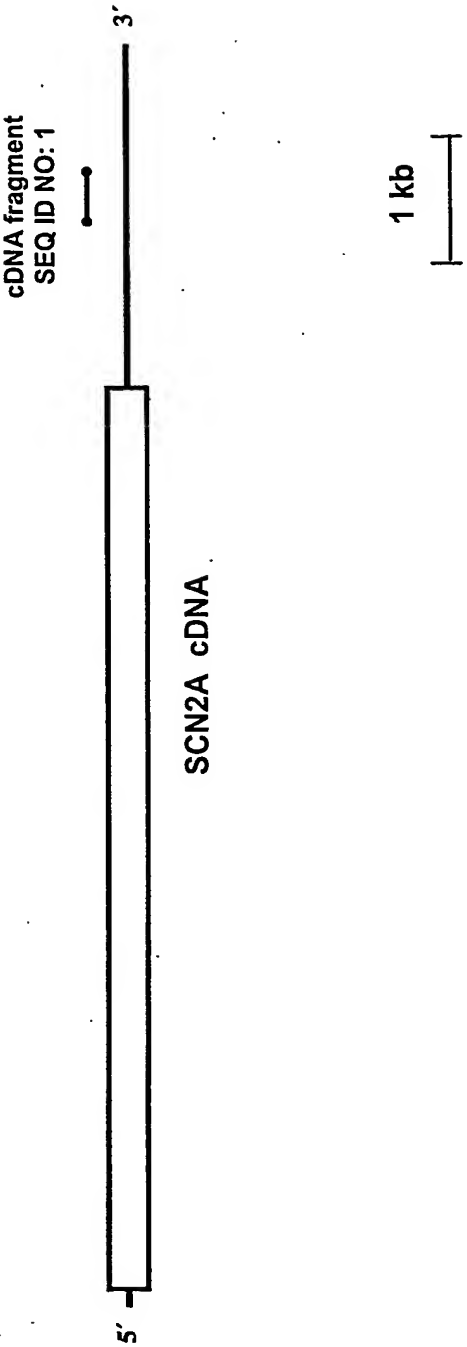


Fig. 7: Alignment of SEQ ID NO: 1 to voltage-gated ion channel type II A cDNA, SEQ ID NO: 2

Length: 272 bp

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272  GTACCCAGTAAGTATAGTTTGGCAAGCTTCCAACAGGTAATATGATGTAA 223
    |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||
7248  GTACCACAGTAAGTATAGTTTGGCAAGCTTCCAACAGGTAATATGATGTAA 7297
    .      .      .      .      .      .      .      .      .      .
222  TTGGTTCCATTATAGTTTGAAGCTGTCACTGCTGCATGTTTATCTTGCCT 173
    |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||
7298  TTGGTTCCATTATAGTTTGAAGCTGTCACTGCTGCATGTTTATCTTGCCT 7347
    .      .      .      .      .      .      .      .      .      .
172  ATGCTGCTGTATCTTATTCCCTTCCCTGTTCAGAAGTCTAATATGGGAAG 123
    |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||
7348  ATGCTGCTGTATCTTATTCCCTTCCCTGTTCAGAAGTCTAATATGGGAAG 7397
    .      .      .      .      .      .      .      .      .      .
122  CCATATATCAGTGGTAAAGTGAAGCAAATTGTTCTACCAAGACCTCATTC 73
    |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||
7398  CCATATATCAGTGGTAAAGTGAAGCAAATTGTTCTACCAAGACCTCATTC 7447
    .      .      .      .      .      .      .      .      .      .
72  TTCATGTCATTAAGCAATAGGTTGCAGCAAACAAGGAAGAGCTTCTTGCT 23
    |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||  |||||
7448  TTCATGTCATTAAGCAATAGGTTGCAGCAAACAAGGAAGAGCTTCTTGCT 7497
    .      .      .      .      .      .      .      .      .      .
22  TTTTATTCTTCCAACCTTAATT 1
    |||||  |||||  |||||  |||||  |||||
7498  TTTTATTCTTCCAACCTTAATT 7519

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**Fig. 8: SEQ ID NO. 2: nucleotide sequence of
human SCN2A cDNA**

Length: 8292 bp

```
1  CACTTTCTTA TGCAAGGAGC TAAACAGTGA TTAAAGGAGC AGGATGAAAA
51  GATGGCACAG TCAGTGCTGG TACCGCCAGG ACCTGACAGC TTCCGCTTCT
101 TTACCAGGGA ATCCCTTGCT GCTATTGAAC AACGCATTGC AGAAGAGAAA
151 GCTAAGAGAC CCAAACAGGA ACGCAAGGAT GAGGATGATG AAAATGGCCC
201 AAAGCCAAAC AGTGACTTGG AAGCAGGAAA ATCTCTTCCA TTTATTTATG
251 GAGACATTCC TCCAGAGATG GTGTCAGTGC CCCTGGAGGA TCTGGACCCC
301 TACTATATCA ATAAGAAAAC GTTTATAGTA TTGAATAAAG GGAAAGCAAT
351 CTCTCGATTG AGTGCCACCC CTGCCCTTTA CATTTTAACT CCCTTCAACC
401 CTATTAGAAA ATTAGCTATT AAGATTTTGG TACATTCCTT ATTCAATATG
451 CTCATTATGT GCACGATTCT TACCAACTGT GTATTTATGA CCATGAGTAA
501 CCCTCCAGAC TGGACAAAGA ATGTGGAGTA TACCTTTACA GGAATTTATA
551 CTTTTGAATC ACTTATTAAA ATACTTGCAA GGGGCTTTTG TTTAGAAGAT
601 TTCACATTTT TACGGGATCC ATGGAATTGG TTGGATTTCA CAGTCATTAC
651 TTTTGCATAT GTGACAGAGT TTGTGGACCT GGGCAATGTC TCAGCGTTGA
701 GAACATTCAG AGTTCTCCGA GCATTGAAAA CAATTCAGT CATTCAGGC
751 CTGAAGACCA TTGTGGGGGC CCTGATCCAG TCAGTGAAGA AGCTTTCTGA
801 TGTCATGATC TTGACTGTGT TCTGTCTAAG CGTGTGTGCG CTAATAGGAT
851 TGCAGTTGTT CATGGGCAAC CTACGAAATA AATGTTTGCA ATGGCCTCCA
901 GATAATTCTT CCTTTGAAAT AAATATCAC T CCTTCTTTA ACAATTCATT
951 GGATGGGAAT GGTACTACTT TCAATAGGAC AGTGAGCATA TTTAACTGGG
1001 ATGAATATAT TGAGGATAAA AGTCACTTTT ATTTTGTAGA GGGGCAAAAT
1051 GATGCTCTGC TTTGTGGCAA CAGCTCAGAT GCAGGCCAGT GTCCTGAAGG
1101 ATACATCTGT GTGAAGGCTG GTAGAAACCC CAACTATGGC TACACGAGCT
1151 TTGACACCTT TAGTTGGGCC TTTTGTCTCT TATTTCGTCT CATGACTCAA
1201 GACTTCTGGG AAAACCTTTA TCAACTGACA CTACGTGCTG CTGGGAAAAC
1251 GTACATGATA TTTTGTGTGC TGGTCATTTT CTTGGGCTCA TTCTATCTAA
1301 TAAATTTGAT CTTGGCTGTG GTGGCCATGG CCTATGAGGA ACAGAATCAG
1351 GCCACATTGG AAGAGGCTGA ACAGAAGGAA GCTGAATTTT AGCAGATGCT
1401 CGAACAGTTG AAAAAGCAAC AAGAAGAAGC TCAGGCGGCA GCTGCAGCCG
1451 CATCTGCTGA ATCAAGAGAC TTCAGTGGTG CTGGTGGGAT AGGAGTTTTT
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1551 GCTGAAAAAC AGAAGAAAGA AAAAGAAACA GAAAGAACAG TCTGGAGAAG
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1701 GAGATTTTCT TCTCCACACC AGTCCTTACT GAGCATCCGT GGCTCCCTTT
1751 TCTCTCCAAG ACGCAACAGT AGGGCGAGCC TTTTCAGCTT CAGAGGTCGA
1801 GCAAAGGACA TTGGCTCTGA GAATGACTTT GCTGATGATG AGCACAGCAC
1851 CTTTGAGGAC AATGACAGCC GAAGAGACTC TCTGTTCTGT CCGCACAGAC
1901 ATGGAGAACG GCGCCACAGC AATGTCAGCC AGGCCAGCCG TGCCTCCAGG
1951 GTGCTCCCCA TCCTGCCCAT GAATGGGAAG ATGCATAGCG CTGTGGACTG
2001 CAATGGTGTG GTCTCCCTGG TCGGGGGCCC TTCTACCCTC ACATCTGCTG
2051 GGCAGCTCCT ACCAGAGGGC ACAACTACTG AAACAGAAAT AAGAAAGAGA
2101 CGGTCCAGTT CTTATCATGT TTCCATGGAT TTATTGGAAG ATCCTACATC
2151 AAGGCAAAGA GCAATGAGTA TAGCCAGTAT TTTGACCAAC ACCATGGAAG
2201 AACTTGAAGA ATCCAGACAG AAATGCCAC CATGCTGGTA TAAATTTGCT
2251 AATATGTGTT TGATTTGGGA CTGTTGTAAA CCATGGTTAA AGGTGAAACA
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2301	CCTTGTC AAC	CTGGTTGTAA	TGGACCCATT	TGTTGACCTG	GCCATCACCA
2351	TCTGCATTGT	CTTAAATACA	CTCTTCATGG	CTATGGAGCA	CTATCCCATG
2401	ACGGAGCAGT	TCAGCAGTGT	ACTGTCTGTT	GGAAACCTGG	TCTTCACAGG
2451	GATCTTCACA	GCAGAAATGT	TTCTCAAGAT	AATTGCCATG	GATCCATATT
2501	ATTACTTTCA	AGAAGGCTGG	AATATTTTTG	ATGGTTTTAT	TGTGAGCCTT
2551	AGTTTAATGG	AACTTGGTTT	GGCAAATGTG	GAAGGATTGT	CAGTTCTCCG
2601	ATCATTCGGG	CTGCTCCGAG	TTTTCAAGTT	GGCAAAATCT	TGGCCAACTC
2651	TAAATATGCT	AATTAAGATC	ATTGGCAATT	CTGTGGGGGC	TCTAGGAAAC
2701	CTCACCTTGG	TATTGGCCAT	CATCGTCTTC	ATTTTTGCTG	TGGTCGGCAT
2751	GCAGCTCTTT	GGTAAGAGCT	ACAAAGAATG	TGTCTGCAAG	ATTTCCAATG
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2851	ATCGTGTTCC	GCGTGCTGTG	TGGAGAGTGG	ATAGAGACCA	TGTGGGACTG
2901	TATGGAGGTC	GCTGGCCAAA	CCATGTGCCT	TACTGTCTTC	ATGATGGTCA
2951	TGGTGATTGG	AAATCTAGTG	GTTCTGAACC	TCTTCTTGGC	CTTGCTTTTTG
3001	AGTTCCTTCA	GTTCTGACAA	TCTTGCTGCC	ACTGATGATG	ATAACGAAAT
3051	GAATAATCTC	CAGATTGCTG	TGGGAAGGAT	GCAGAAAGGA	ATCGATTTTG
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3151	AAAGCTTTAG	ATGAAATTAA	ACCGCTTGAA	GATCTAAATA	ATAAAAAAGA
3201	CAGCTGTATT	TCCAACCATA	CCACCATAGA	AATAGGCAAA	GACCTCAATT
3251	ATCTCAAAGA	CGGAAATGGA	ACTACTAGTG	GCATAGGCAG	CAGTGTAGAA
3301	AAATATGTCT	TGGATGAAAG	TGATTACATG	TCATTTATAA	ACAACCCTAG
3351	CCTCACTGTG	ACAGTACCAA	TTGCTGTTGG	AGAATCTGAC	TTTGAAAATT
3401	TAAATACTGA	AGAATTTCAGC	AGCGAGTCAG	ATATGGAGGA	AAGCAAAGAG
3451	AAGCTAAATG	CAACTAGTTC	ATCTGAAGGC	AGCACGGTTG	ATATTGGAGC
3501	TCCCGCCGAG	GGAGAACAGC	CTGAGGTTGA	ACCTGAGGAA	TCCCTTGAAC
3551	CTGAAGCCTG	TTTTACAGAA	GACTGTGTAC	GGAAGTTCAA	GTGTTGTCAG
3601	ATAAGCATAG	AAGAAGGCAA	AGGGAAACTC	TGGTGGAATT	TGAGGAAAAC
3651	ATGCTATAAG	ATAGTGGAGC	ACAATTGGTT	CGAAACCTTC	ATTGTCTTCA
3701	TGATTCTGCT	GAGCAGTGGG	GCTCTGGCCT	TTGAAGATAT	ATACATTGAG
3751	CAGCGAAAAA	CCATTAAGAC	CATGTTAGAA	TATGCTGACA	AGGTTTTTCA
3801	TTACATATTC	ATTCTGGAAA	TGCTGCTAAA	GTGGGTTGCA	TATGGTTTTT
3851	AAGTGTATTT	TACCAATGCC	TGGTGCTGGC	TAGACTTCCT	GATTGTTGAT
3901	GTCTCACTGG	TTAGCTTAAC	TGCAAATGCC	TTGGGTTACT	CAGAACTTGG
3951	TGCCATCAAA	TCCCTCAGAA	CACTAAGAGC	TCTGAGGCCA	CTGAGACTTT
4001	TGTCCCGGTT	TGAAGGAATG	AGGGTTGTTG	TAAATGCTCT	TTTAGGAGCC
4051	ATTCCATCTA	TCATGAATGT	ACTTCTGGTT	TGTCTGATCT	TTTGCTAAT
4101	ATTCACTATC	ATGGGAGTGA	ATCTCTTTGC	TGGCAAGTTT	TACCATTGTA
4151	TTAATTACAC	CACTGGAGAG	ATGTTTGATG	TAAGCGTGGT	CAACAACCTAC
4201	AGTGAGTGCA	AAGCTCTCAT	TGAGAGCAAT	CAAACCTGCC	GGTGGA AAAA
4251	TGTGAAAGTA	AACTTTGATA	ACGTAGGACT	TGGATATCTG	TCTCTACTTC
4301	AAGTAGCCAC	GTTTAAGGGA	TGGATGGATA	TTATGTATGC	AGCTGTTGAT
4351	TCACGAAATG	TAGAATTACA	ACCCAAGTAT	GAAGACAACC	TGTACATGTA
4401	TCTTTATTTT	GTCATCTTTA	TTATTTTTTG	TTCATTCTTT	ACCTTGAATC
4451	TTTTTCATTG	TGTCATCATA	GATAACTTCA	ACCAACAGAA	AAAGAAGTTT
4501	GGAGGTCAAG	ACATTTTTTAT	GACAGAAGAA	CAGAAGAAAT	ACTACAATGC
4551	AATGAAAAAA	CTGGGTTCAA	AGAAACCACA	AAAACCCATA	CCTCGACCTG
4601	CTAACAAATT	CCAAGGAATG	GTCTTTGATT	TTGTAACCAA	ACAAGTCTTT
4651	GATATCAGCA	TCATGATCCT	CATCTGCCTT	AACATGGTCA	CCATGATGGT
4701	GGAAACCGAT	GACCAGAGTC	AAGAAATGAC	AAACATTCTG	TACTGGATTA
4751	ATCTGGTGTT	TATTGTTCTG	TTCAGTGGAG	AATGTGTGCT	GAAACTGATC
4801	TCTCTTCGTT	ACTACTATTT	CACTATTGGA	TGGAATATTT	TTGATTTTGT
4851	GGTGGTCATT	CTCTCCATTG	TAGGAATGTT	TCTGGCTGAA	CTGATAGAAA
4901	AGTATTTTTG	GTCCCCTACC	CTGTTCCGAG	TGATCCGTCT	TGCCAGGATT
4951	GGCCGAATCC	TACGWCTGAT	CAAAGGAGCA	AAGGGGATCC	GCACGCTGCT
5001	CTTTGCTTTG	ATGATGTCCC	TTCTGCGTT	GTTTAACATC	GGCCTCCTTC
5051	TTTTCTGGT	CATGTTTCATC	TACGCCATCT	TTGGGATGTC	CAATTTTGCC
5101	TATGTTAAGA	GGGAAGTTGG	GATCGATGAC	ATGTTCAACT	TTGAGACCTT

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5201	ATGGATTGCT	AGCACCTATT	CTTAATAGTG	GACCTCCAGA	CTGTGACCCT
5251	GACAAAGATC	ACCTTGGAAG	CTCAGTTAAA	GGAGACTGTG	GGAACCCATC
5301	TGTTGGGATT	TTCTTTTTTG	TCAGTTACAT	CATCATATCC	TTCTTGTTG
5351	TGGTGAACAT	GTACATCGCG	GTCATCCTGG	AGAACTTCAG	TGTTGCTACT
5401	GAAGAAAGTG	CAGAGCCTCT	GAGTGAGGAT	GACTTTGAGA	TGTTCTATGA
5451	GGTTTGGGAG	AAGTTTGATC	CCGATGCGAC	CCAGTTTATA	GAGTTTGCCA
5501	AACTTTCTGA	TTTTGCAGAT	GCCCTGGATC	CTCCTCTTCT	CATAGCAAAA
5551	CCCAACAAAG	TCCAGCTCAT	TGCCATGGAT	CTGCCCATGG	TGAGTGGTGA
5601	CCGGATCCAC	TGTCTTGACA	TCTTATTTGC	TTTTACAAAG	CGTGTTTTGG
5651	GTGAGAGTGG	AGAGATGGAT	GCCCTTCGAA	TACAGATGGA	AGAGCGATTCT
5701	ATGGCATCAA	ACCCCTCCAA	AGTCTCTTAT	GAGCCCATTA	CGACCACGTT
5751	GAAACGCAAA	CAAGAGGAGG	TGTCTGCTAT	TATTATCCAG	AGGGCTTACA
5801	GACGCTACCT	CTTGAAGCAA	AAAGTTAAAA	AGGTATCAAG	TATATACAAG
5851	AAAGACAAAG	GCAAAGAATG	TGATGGAACA	CCCATCAAAG	AAGATACTCT
5901	CATTGATAAA	CTGAATGAGA	ATTCAACTCC	AGAGAAAACC	GATATGACGC
5951	CTTCCACCAC	GTCTCCACCC	TCGTATGATA	GTGTGACCAA	ACCAGAAAAA
6001	GAAAAATTTG	AAAAAGACAA	ATCAGAAAAAG	GAAGACAAAG	GGAAAGATAT
6051	CAGGGAAAGT	AAAAAGTAAA	AAGAAACCAA	GAATTTTCCA	TTTTGTGATC
6101	AATTGTTTAC	AGCCCGTGAT	GGTGATGTGT	TTGTGTCAAC	AGGACTCCCA
6151	CAGGAGGTCT	ATGCCAAACT	GACTGTTTTT	ACAAATGTAT	ACTTAAGGTC
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6251	AACTGGAGAA	ATAGTATCGA	TGGGAGGTTT	CTATTTTCAC	AACCAGCTGA
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6351	TTAAAGGGGG	GAGGGAAGTT	AAATTTTTAT	GTAAATTCAA	CATGTGACAC
6401	TTGATAATAG	TAATTGTCAC	CAGTGTTTAT	GTTTTAACTG	CCACACCTGC
6451	CATATTTTTA	CAAAACGTGT	GCTGTGAATT	TATCACTTTT	CTTTTTAATT
6501	CACAGGTTGT	TTACTATTAT	ATGTGACTAT	TTTTGTAAAT	GGGTTTGTGT
6551	TTGGGGAGAG	GGATTAAAGG	GAGGGAATTC	TACATTTCTC	TATTGTATTG
6601	TATAACTGGA	TATATTTTAA	ATGGAGGCAT	GCTGCAATTC	TCATTCACAC
6651	ATAAAAAAAT	CACATCACAA	AAGGGAAGAG	TTTACTTCTT	GTTTCAGGAT
6701	GTTTTTAGAT	TTTGAGGTG	CTTAAATAGC	TATTCGTATT	TTAAGGGTGT
6751	CTCATCCAGA	AAAAATTTAA	TGTGCCTGTA	AATGTTCCAT	AGAATCACAA
6801	GCATTAAAGA	GTTGTTTTAT	TTTTACATAA	CCCATTAAAT	GTACATGTAT
6851	ATATGTATAT	ATGTATATGT	CGGTGTATAT	ACATATATAT	GTATACACAC
6901	ATGCACACAC	AGAGATATAC	ACATACCATT	ACATTGTCAT	TCACAGTCCC
6951	AGCAGCATGA	CTATCACATT	TTTGATAAGT	GTCCTTTGGC	ATAAAATAAA
7001	AATATCCTAT	CAGTCCTTTC	TAAGAAGCCT	GAATTGACCA	AAAAACATCC
7051	CCACCACCAC	TTTATAAAGT	TGATTCTGCT	TTATCCTGCA	GTATTGTTTA
7101	GCCATCTTCT	GCTCTTGGTA	AGGTTGACAT	AGTATATGTC	AATTTAAAAA
7151	ATAAAAGTCT	GCTTTGTAAA	TAGTAATTTT	ACCCAGTGGT	GCATGTTTGA
7201	GCAAACAAAA	ATGATGATTT	AAGCACACTA	CTTATTGCAT	CAAATATGTA
7251	CCACAGTAAG	TATAGTTTGC	AAGCTTTCAA	CAGGTAATAT	GATGTAATTG
7301	GTTCCATTAT	AGTTTGAAGC	TGTCACTGCT	GCATGTTTAT	CTTGCCTATG
7351	CTGCTGTATC	TTATTCCTTC	CACTGTTCAG	AAGTCTAATA	TGGGAAGCCA
7401	TATATCAGTG	GTAAAGTGAA	GCAAATTGTT	CTACCAAGAC	CTCATTCTTC
7451	ATGTCATTAA	GCAATAGGTT	GCAGCAAACA	AGGAAGAGCT	TCTTGCTTTT
7501	TATTCTTCCA	ACCTTAATTG	AACACTCAAT	GATGAAAAGC	CCGACTGTAC
7551	AAACATGTTG	CAAGCTGCTT	AAATCTGTTT	AAAATATATG	GTTAGAGTTT
7601	TCTAAGAAAA	TATAAATACT	GTAATAAAGT	CATTTTATTT	TATTTTTTCA
7651	CCTTTTGTAC	GTAAAATGAG	AAATTAAGAAG	TATCTTCAGG	TGGATGTCAC
7701	AGTCACTATT	GTTAGTTTCT	GTTCTTAGCA	CTTTTAAATT	GAAGCACTTC
7751	ACAAAATAAG	AAGCAAGGAC	TAGGATGCAG	TGTAGGTTTC	TGCTTTTTTA
7801	TTAGTACTGT	AAACTTGCAC	ACATTTCAAT	GTGAAACAAA	TCTCAAACCTG
7851	AGTTCAATGT	TTATTTGCTT	TCAATAGTAA	TGCCTTATCA	TTGAAAGAGG
7901	CTTAAAGAAA	AAAAAATCA	GCTGATACTC	TTGGCATTGC	TTGAATCCAA
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11/13

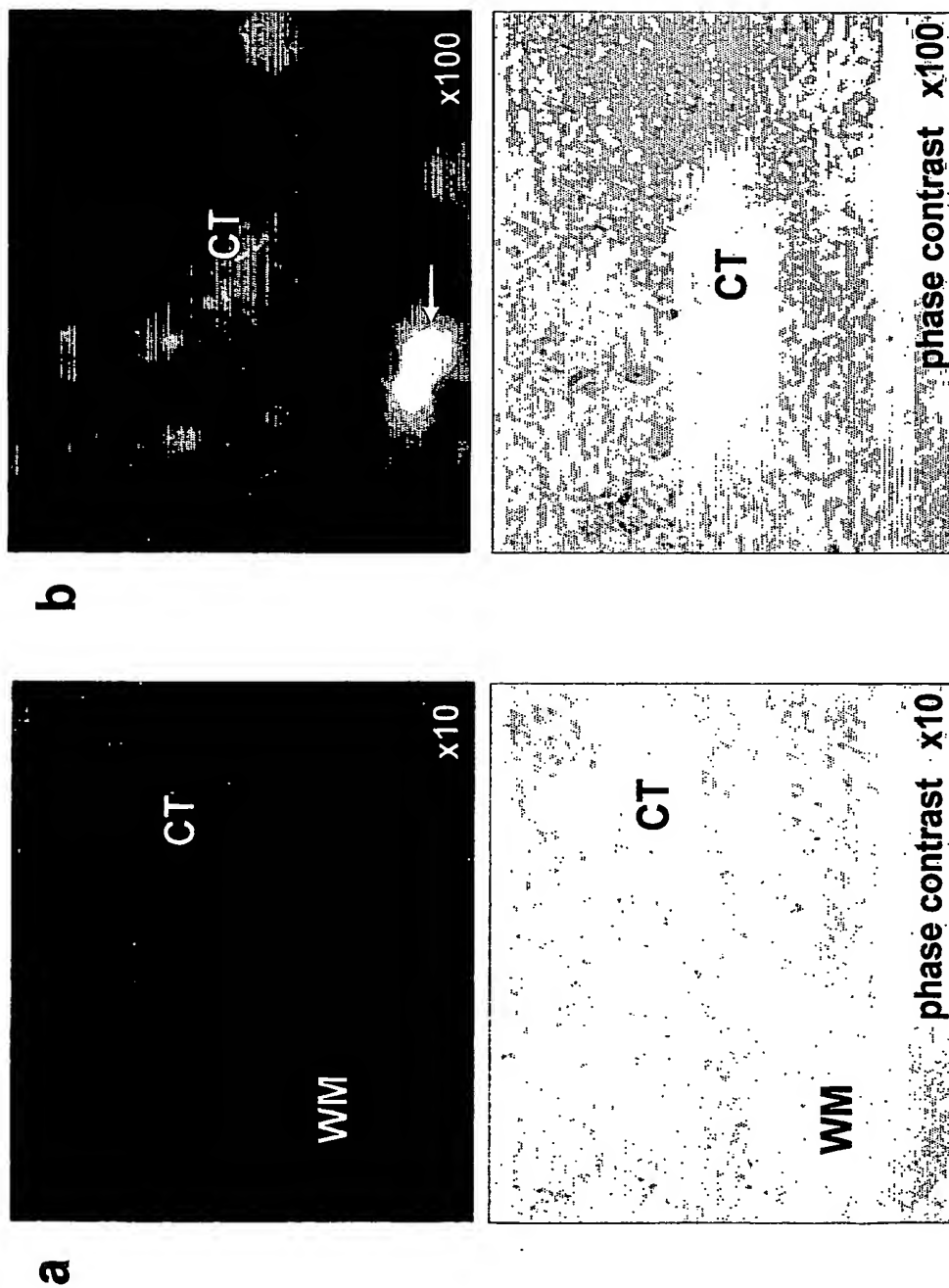
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8151	AATAAATGTA	GATTCTTTAT	ACTGAAGCTA	TTGACTTGTA	GTGTGTTGGT
8201	GAAATGCATG	CAGGAAAATG	CTGTTACCAT	AAAGAACGGT	AAACCACATT
8251	ACAATCAAGC	CAAAAGAATA	AAGGTTTCGC	TTTTGTTTTT	GT

Fig. 9: SEQ ID NO. 3:
amino acid sequence of
human SCN2A protein

Length: 2005 aa

1	MAQSVLVPPG	PDSFRFFTRE	SLAAIEQRIA	EEKAKRPKQE	RKDEDDENG
51	KPNSDLEAGK	SLPFIYGDIP	PEMVSVPLED	LDPYIINKKT	FIVLNKGKAI
101	SRFSATPALY	ILTPFNPIRK	LAIKILVHSL	FNMLIMCTIL	TNCVFM TMSN
151	PPDWTKNVEY	TFTGIYTFES	LIKILARGFC	LEDFTFLRDP	WNWLDFTVIT
201	FAYVTEFVDL	GNVSALRTR	VLRALKTISV	IPGLKTI VGA	LIQSVKKLSD
251	VMILTVFCLS	VFALIGLQLF	MGNLRNKCLQ	WPPDNSSFEI	NITSFFNNSL
301	DGNGTTFNRT	VSIFNWDEYI	EDKSHFYFLE	GQNDALLCGN	SSDAGQCPEG
351	YICVKAGRNP	NYGYTSFDTF	SWAFLSLFRL	MTQDFWENLY	QLTLRAAGKT
401	YMIFFVLVIF	LGSFYLINLI	LAVVAMAYEE	QNQATLEEAE	QKEAEFQOML
451	EQLKKQQEEA	QAAAAAASAE	SRDFSGAGGI	GVFSESSSVA	SKLSSKSEKE
501	LKNRRKKKKQ	KEQSGEEKN	DRVLKSESED	SIRRKGFRRS	LEGSRLTYEK
551	RFSSPHQSL	SIRGSLFSR	RNSRASLFSF	RGRAKDIGSE	NDFADDEHST
601	FEDNDSRRDS	LFVPHRHGER	RHSNVSQASR	ASRVLPILPM	NGKMMSAVDC
651	NGVVSLVGGP	STLTSAGQLL	PEGTTTETEI	RKRRSSSYHV	SMDLLEDPTS
701	RQRAMSIASI	LTNTMEELEE	SRQKCPPCWY	KFANMCLIWD	CCKPWLKVKH
751	LVNLVMDPF	VDLAITICIV	LNTLFMAMEH	YPMTEQFSSV	LSVGNLVFTG
801	IFTAEMFLKI	IAMPYYYFQ	EGWNIFDGF	VSLSLMELGL	ANVEGLSVLR
851	SFRLLRVFKL	AKSWPTLNML	IKIIGNSVGA	LGNLTLVLAI	IVFIFAVVGM
901	QLFGKSYKEC	VCKISNDCEL	PRWHMHDFH	SFLIVFRVLC	GEWIETMWDC
951	MEVAGQTMCL	TVFMMVMVIG	NLVVLNLF	LLSSSFSSDN	LAATDDDNEM
1001	NNLQIAVGRM	QKGIDFVKRK	IREFIQKAFV	RKQKALDEIK	PLEDLNNKKD
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1101	LTVTVPIAVG	ESDFENLNT	EFSSSEDMEE	SKEKLNATSS	SEGSTVDIGA
1151	PAEGEQPEVE	PEESLEPEAC	FTEDCVRKFK	CCQISIEEGK	GKLWWNL RKT
1201	CYKIVEHNWF	ETFIVFMILL	SSGALAFEDI	YIEQRKTIKT	MLEYADKVFT
1251	YIFILEMLLK	WVAYGFQVYF	TNAWCWLDL	IVDVSLVSLT	ANALGYSELG
1301	AIKSLRTRLA	LRPLRALS	EGMRAVVNAL	LGAIPSIMNV	LLVCLIFWLI
1351	FSIMGVNLFA	GKFYHCINYT	TGEMFDVSVV	NNYSECKALI	ESNQ TARWKN
1401	VKVNFDNVGL	GYL SLLQVAT	FKGWM DIMYA	AVDSRNVELQ	PKYEDNLYMY
1451	LYFVIFIIIFG	SFFT LNLFIG	VIIDNFNQOK	KKFGGQDIFM	TEEQKKYYNA
1501	MKKLGSKKPQ	KPIPRPANKF	QGMVDFVTK	QVFDISIMIL	ICLNMVTMMV
1551	ETDDQSQEMT	NILYWINLVF	IVLFTGECVL	KLISLRYYYF	TIGWNIFDFV
1601	VVILSIVGMF	LAELIEKYFV	SPTLFRVIRL	ARIGRILRLI	KGAKGIRTL
1651	FALMMSLPAL	FNIGLL LFLV	MFIYAIFGMS	NFAYVKREVG	IDDMFNFETF
1701	QNSMICLFQI	TTSAGWDGLL	APILN SGPPD	CDPKDHPGS	SVKGD CGNPS
1751	VGIFF FVS	II SFLVVLNM	YIAVILENFS	VATEESAEP	SEDDFEMFYE
1801	VWEKFDPDAT	QFIEFAKLSD	FADALDPPLL	IAKPNKVQLI	AMDLP MVSGD
1851	RIHCLDILFA	FTKRVLGESG	EMDALRIQME	ERFMASNP	VSYPITTTL
1901	KRKQEEVSAI	IIQRAYRRYL	LKQKVKKVSS	IYKKDKGKEC	DGTPIKEDTL
1951	IDKLNENSTP	EKTDMPSTT	SPPSYDSVTK	PEKEKFEKDK	SEKEDKGKDI
2005	RESKK				

**Fig. 10: Images of the human cerebral cortex
labeled with anti SCN2A antibody and with DAPI**



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